

FIG. 1

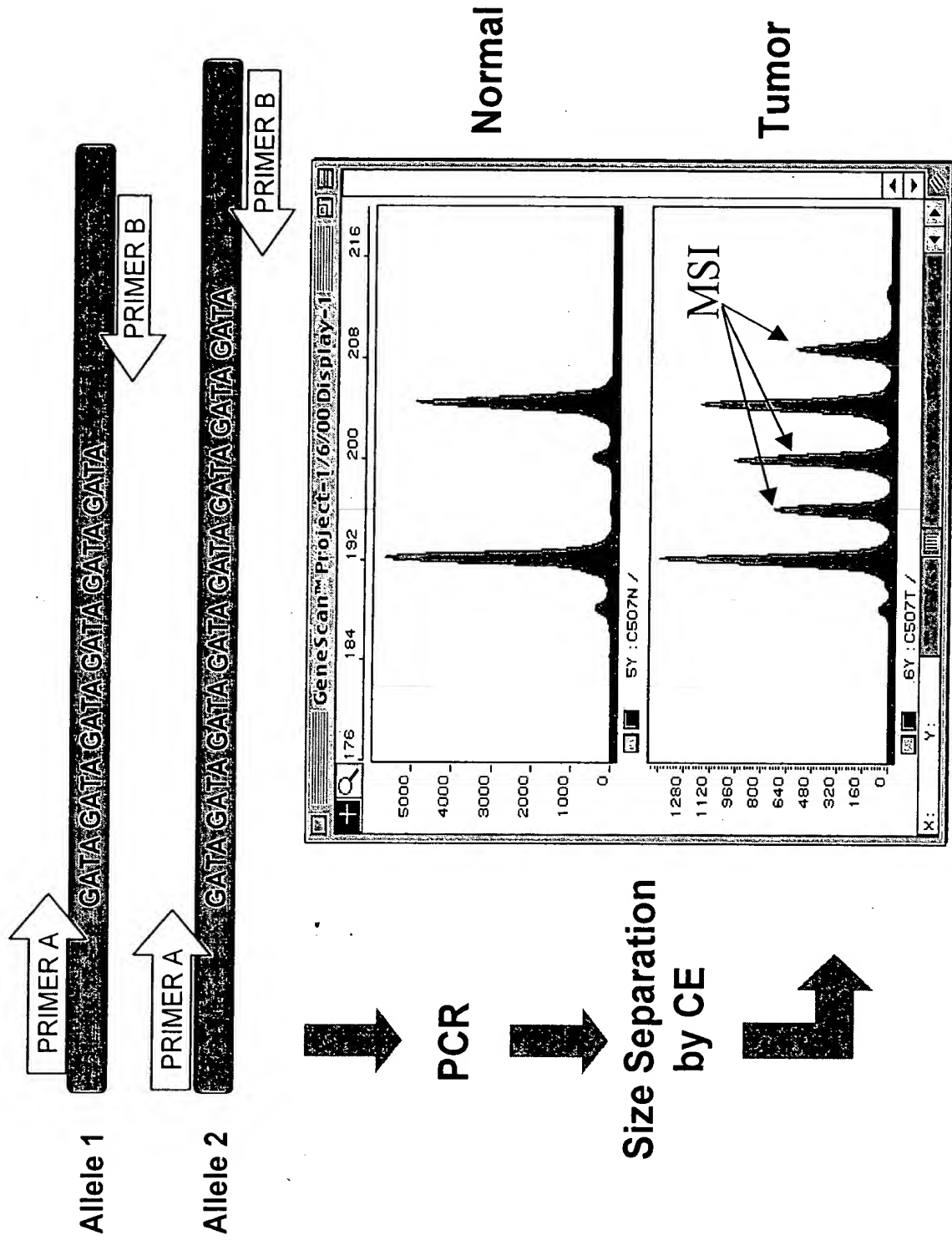
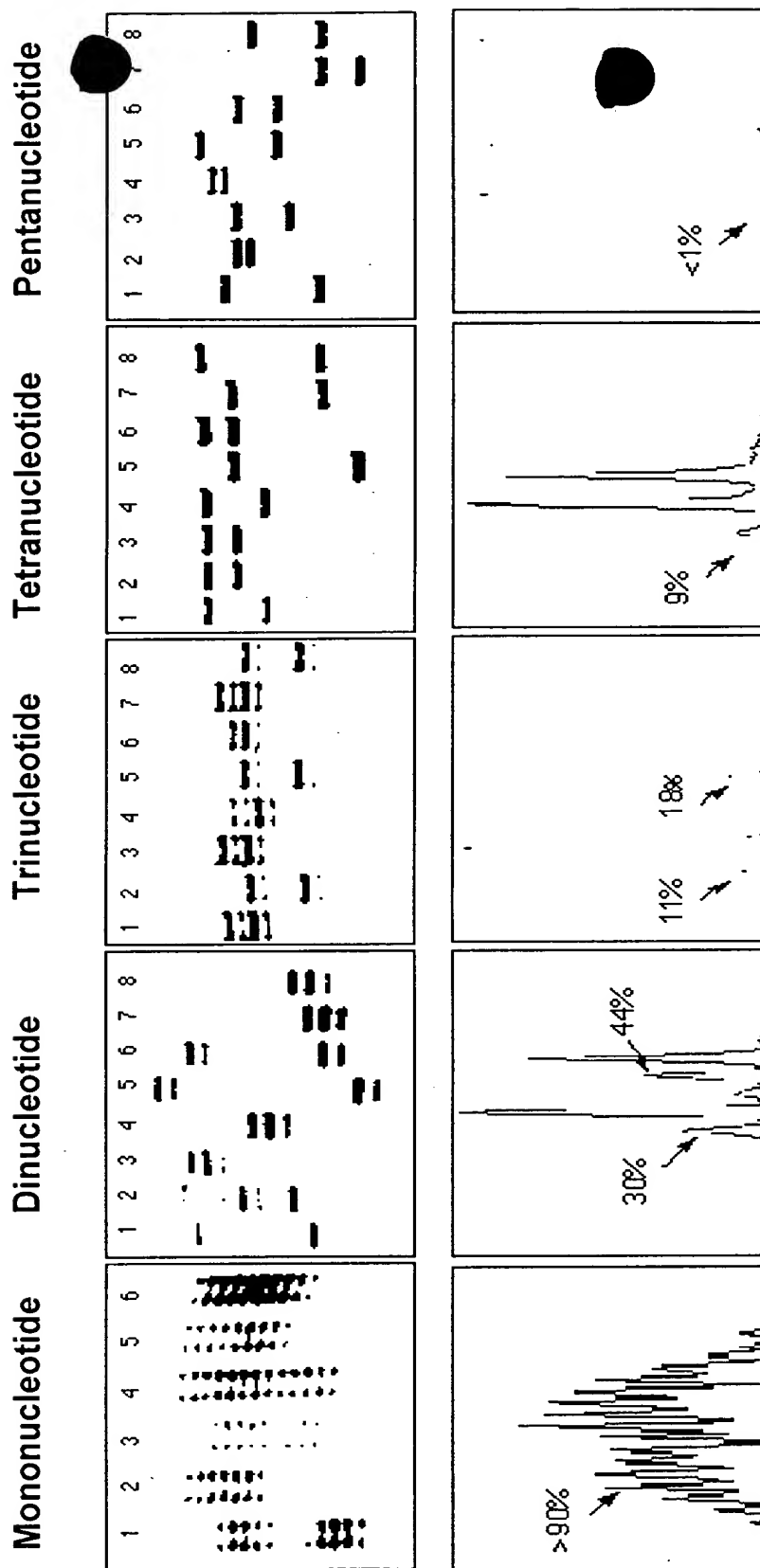


FIG. 2



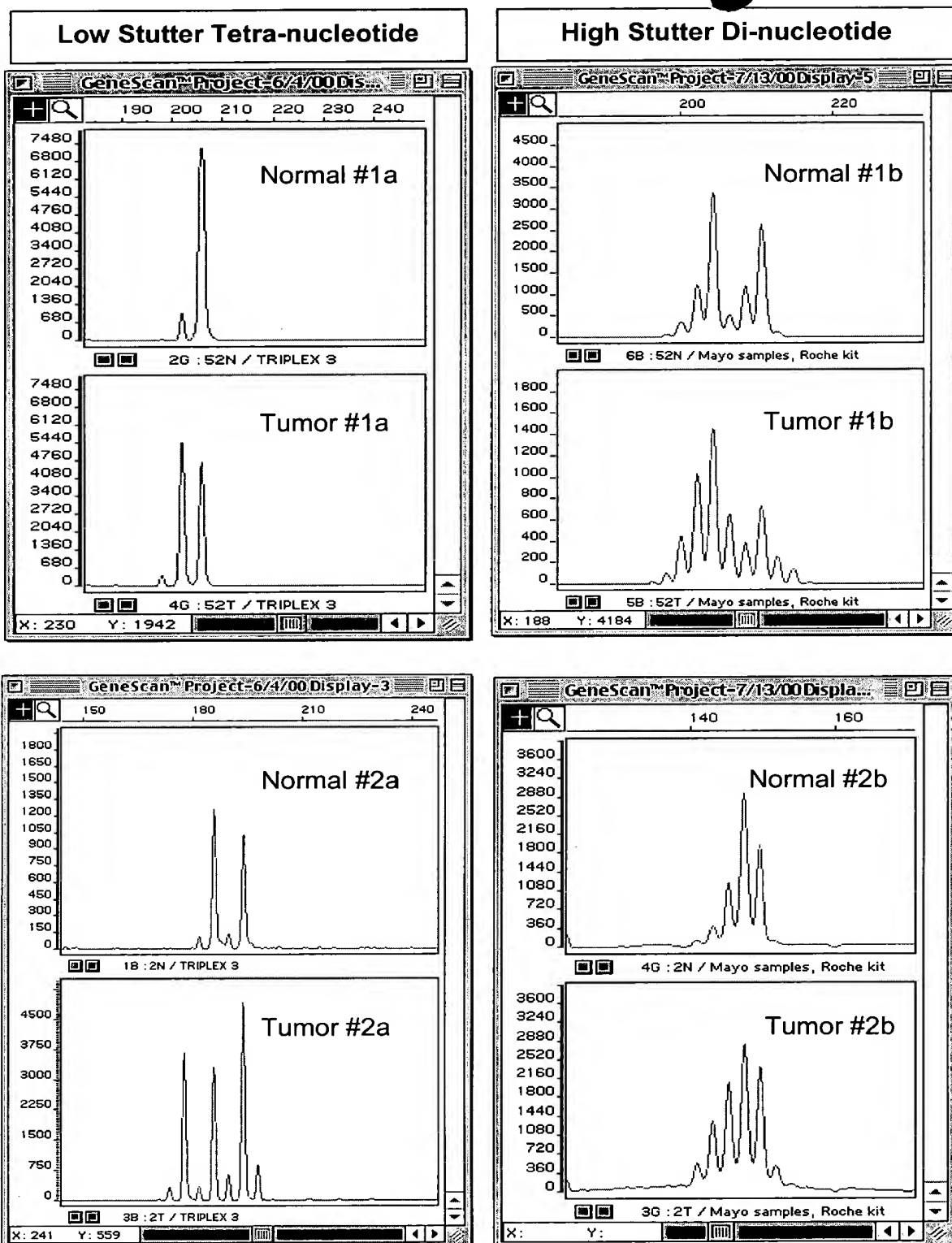


FIG. 3

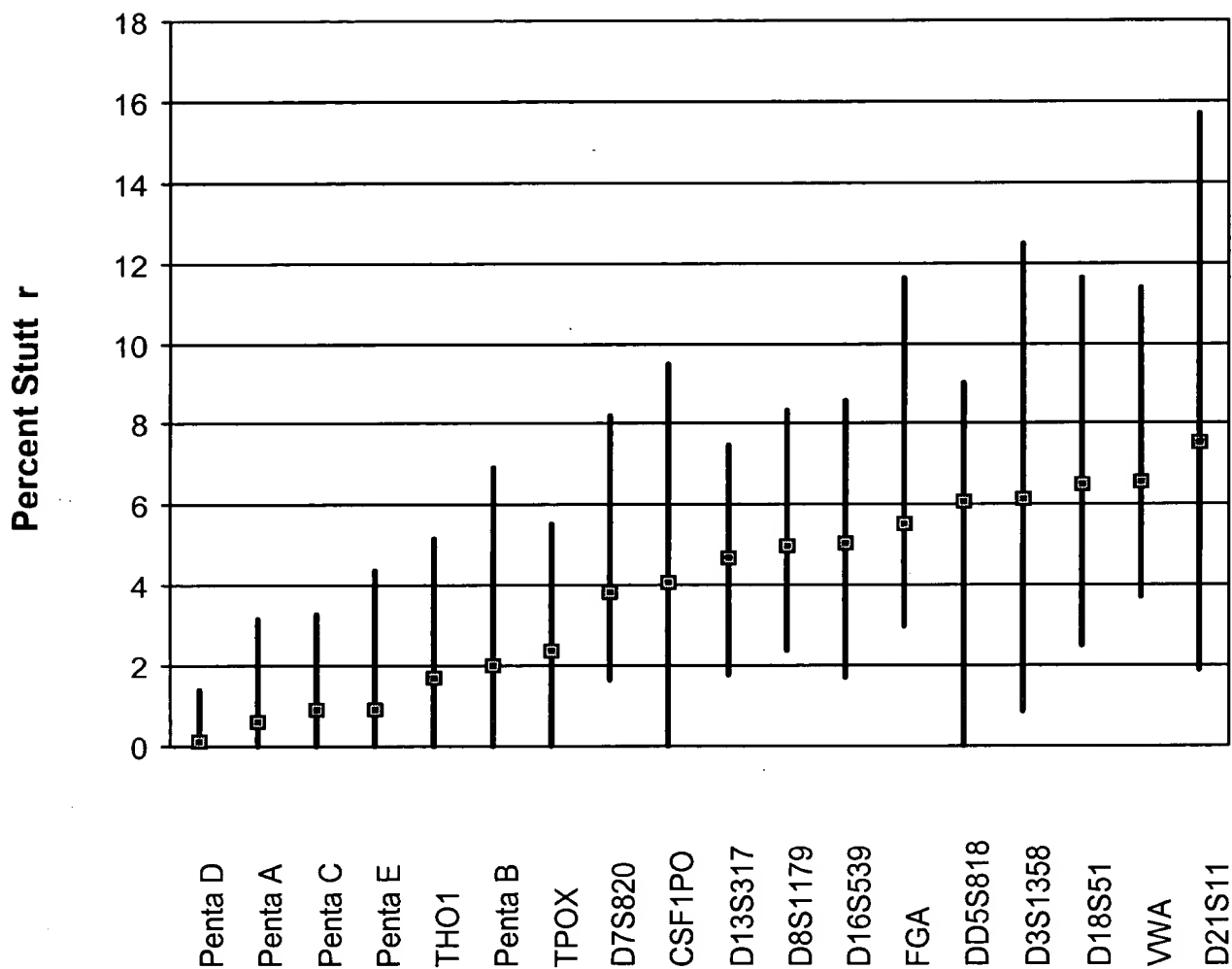


FIG. 4

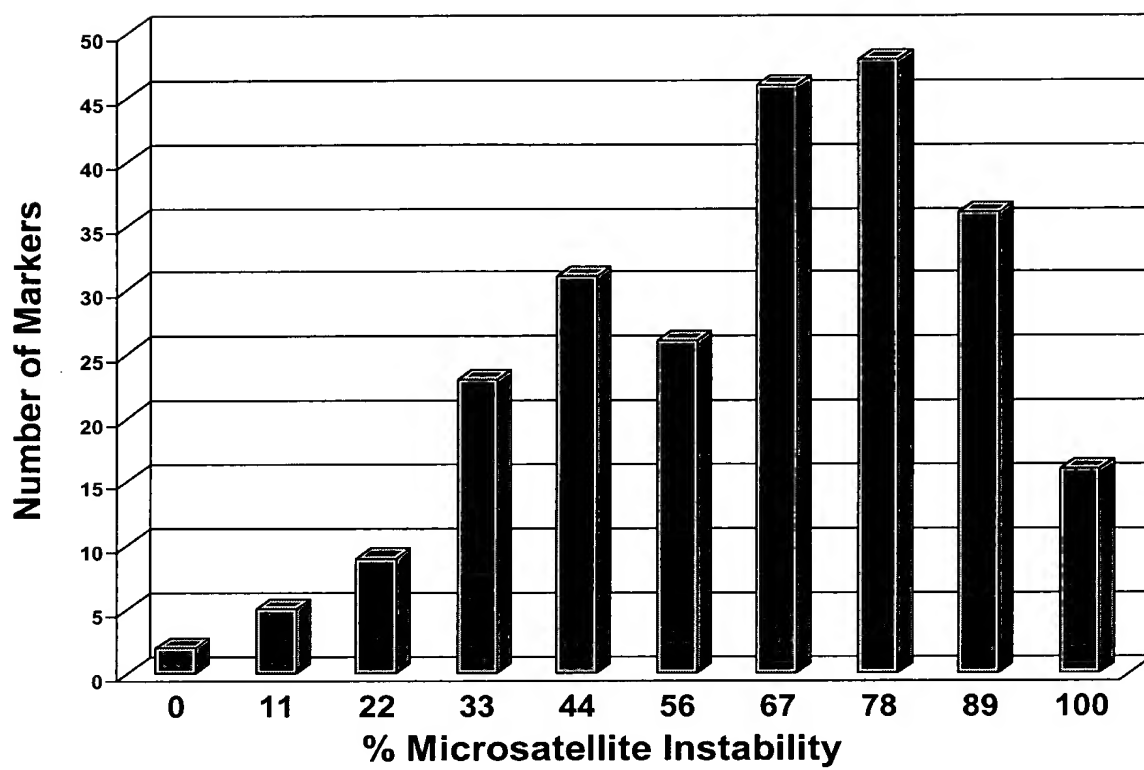


FIG. 5

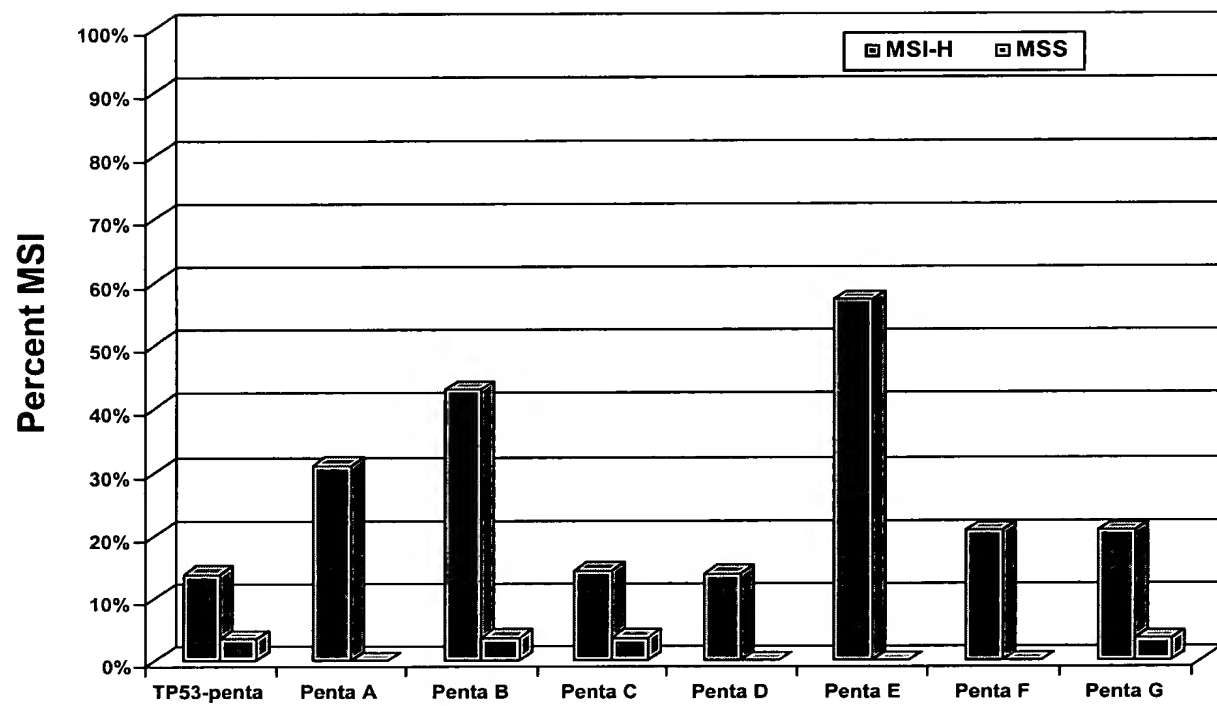


FIG. 6

Figure 1 displays two GeneScan™ Project-6/4/00Display-1 and Display-4, showing electropherogram traces for four samples (A, B, C, D) under Normal and Tumor conditions. The traces are arranged in two columns, with Sample A and B on the left, and Sample C and D on the right. Each sample has two traces: Normal (top) and Tumor (bottom). The traces show peaks corresponding to the samples, with the Tumor traces showing a distinct shift or change in peak intensity compared to the Normal traces. The x-axis represents time or position, with markers at 90 and 120. The y-axis represents signal intensity, with scales ranging from 0 to 3000 for Normal traces and 0 to 1500 for Tumor traces. The bottom of each display shows coordinates X: 98, Y: 103 for Display-1 and X: 104, Y: 707 for Display-4.

FIG. 7

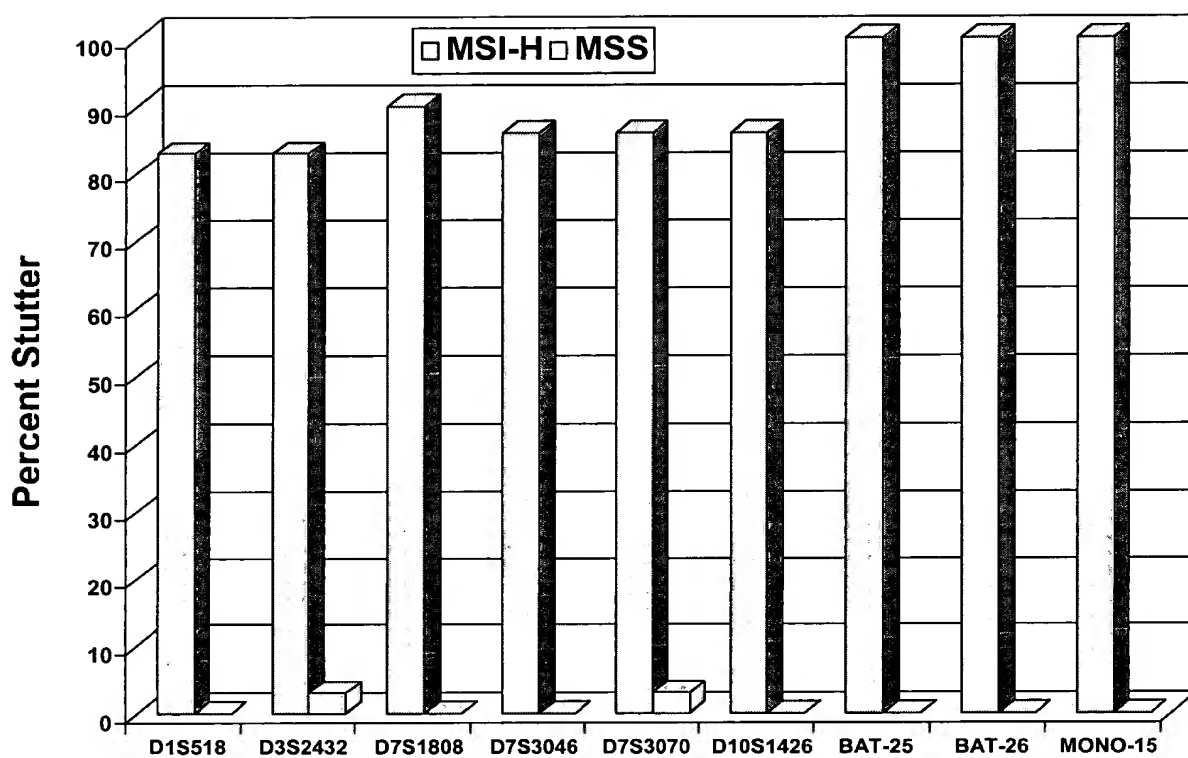


FIG. 8



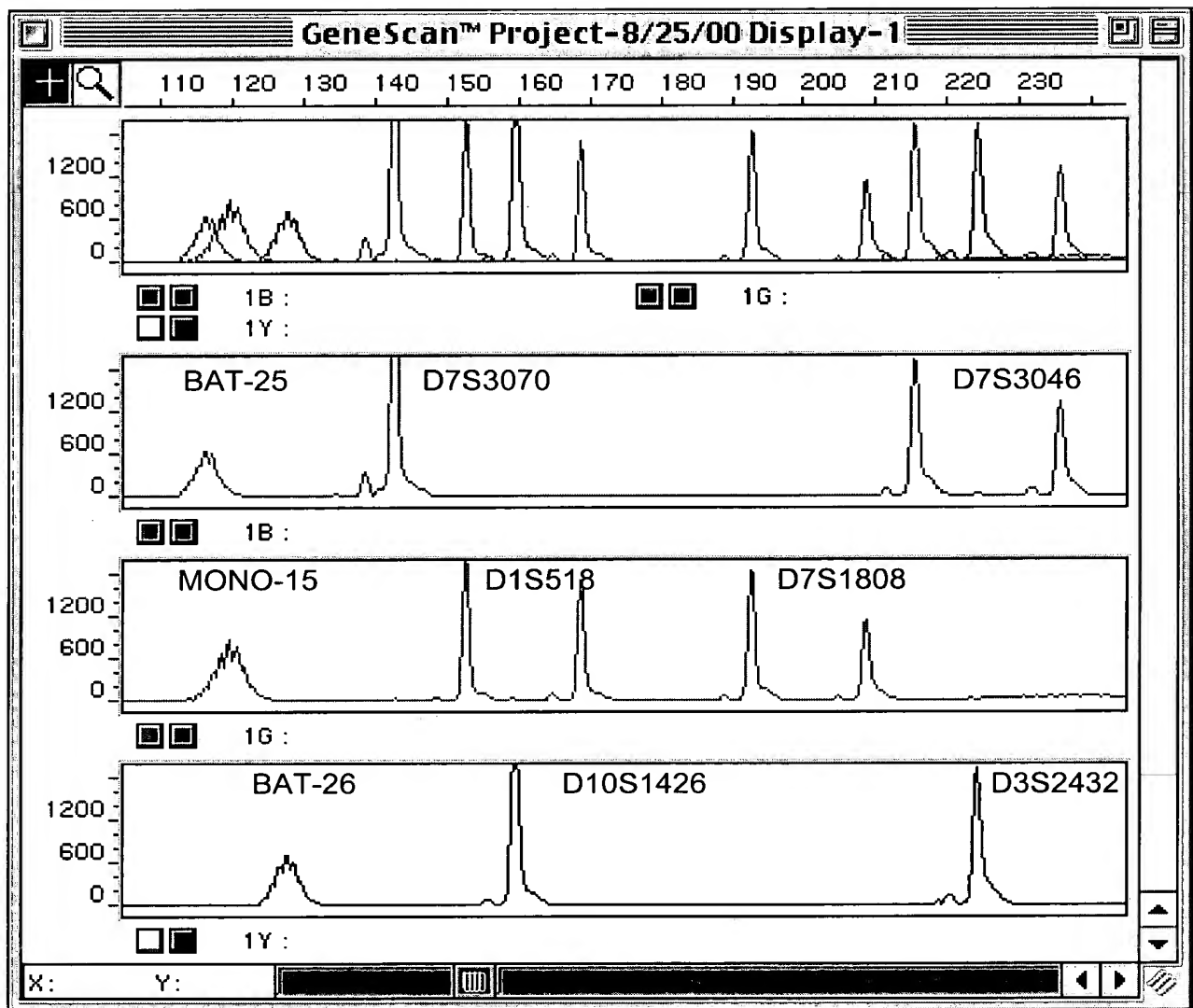


FIG. 9

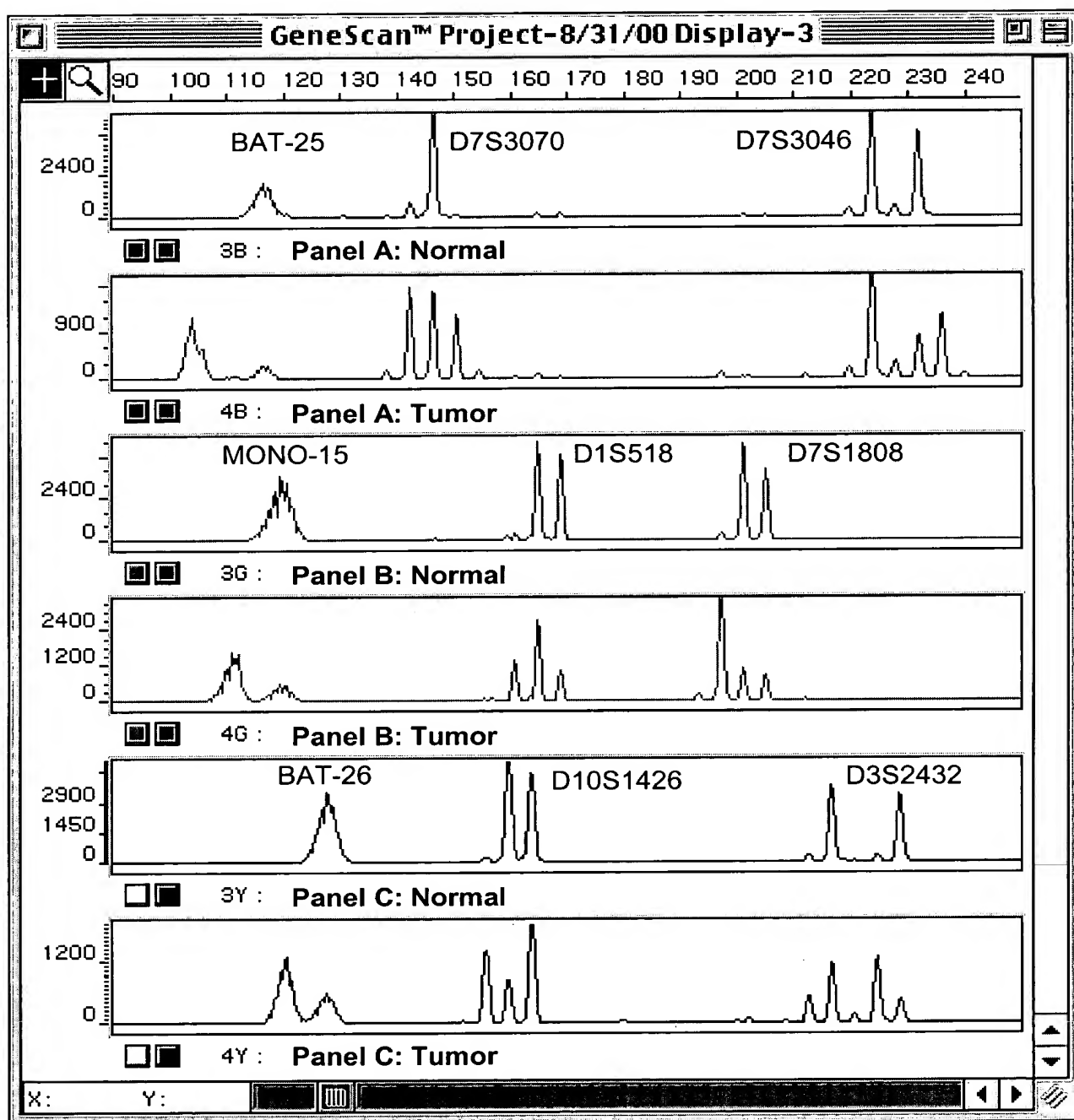


FIG. 10

GeneScan™ Project-8/31/00 Display-4

90 100 110 120 130 140 150 160 170 180 190 200 210 220 230 240

BAT-25 D7S3070 D7S3046

11B : Panel A: Normal

560

12B : Panel A: Tumor

3000

MONO-15 D1S518 D7S1808

11G : Panel B: Normal

600

12G : Panel B: Tumor

1680

BAT-26 D10S1426 D3S2432

11Y : Panel C: Normal

400

12Y : Panel C: Tumor

X: Y:

FIG. 11

GeneScan™ Project-8/31/00 Display-1

100 110 120 130 140 150 160 170 180 190 200 210 220 230

BAT-25 D7S3070 D7S3046

18B : Panel A: Normal

17B : Panel A: Tumor

MONO-15 D1S518 D7S1808

18G : Panel B: Normal

17G : Panel B: Tumor

BAT-26 D10S1426 D3S2432

18Y : Panel C: Normal

17Y : Panel C: Tumor

X: 220 Y: 317

3

